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AX049980.1 GI:12226357 synthetic construct. synthetic construct artificial sequence. 1 (bases 1 to 27) Bout, A., Havenga, M.J. and Vogels, R. Adenovirus derived gene delivery vehicles comprising at leas element of adenovirus type 35 Patent: WO 0070071-A 81 23-NOV-2000; Introgene B.V. (NL) Location/Qualifiers 1. 27 1. 27 /organism="synthetic construct" /db_xref="taxon:32630"	.1	ALIGNMENTS	ωωι	υω.	, w w	ωω.	ω.	4.	4.	4.	4.	5.5	·	55.0		л (л		155.	л .	. ·	100	ν Oυ	5.0		۰.9	9.
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Unclassified.
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1 (bases 1 to 30)
2 1 (bases 1 to 30)
S Kumagai,M.H. and Sverlow,G.G.
Pichia pastoris alcohol oxidase ZZA1 and ZZ:
for heterologous gene expression
Patent: US 5641661-A 8 24-JUN-1997;
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GTGGAAGGCAGTGGAATTTCT 6
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Derose, R. and Hoffmann, J.
GENE CODING FOR ANDROCTONINE, VECTOR CONTAINING
TRANSFORMED DISEASE-RESISTANT PLANTS OBTAINED
Patent: WO 9909189-A 6 25-FEB-1999;
DEROSE RICHARD (FR); HOFFMANN JULES (FR)
LOCATION/QUALIFIERS
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Eukaryota; Metazoa;
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Chenchik,A., Jokhadze,G. and Bibilashvilli,R.
Methods of assaying differential expression
Patent: US 5994076-A 311 30-NOV 1999;
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/db_xref="taxon:9606"
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Sequence 51
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1 (bases 1 to 48)
Bishop, D.H.L. and Kang, C.Y.
Expression of hepatitis b viral antigens
baculovirus vectors
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ent: US 5989538-A 51 23-NOV-1999;
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synthetic construct
artificial sequence.
1 (bases 1 to 48)
Bishop,D.H.L. and Kang,C.Y.
Expression of hepatitis b viral antigens
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A06323
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Bishop,
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Asteridae; euasterids I; Gentianales; Apocynaceae; Rauvolfioideae; Vinceae; Catharanthus.

1 (bases 1 to 48)
                                                         Catharanthus roseus
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16; Conserv
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16; Conserv
                                                                     Madagascar periwinkle.
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David H.L.; Kang, Chil-Yong
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/db_xref="taxon:32630"
12 c 14 g 11 t
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/protein_id="CAA00558.1"
/protein_id="CAA00558.1"
/db_xref="G1:412824"
/translation="MQWNSTALHQALQ"
/translation="12 g 11 t
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A59036
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Scheiflinger, F.D., Antoine, G.D., Falkner, Falko-Guenter, D.,
Dorner, F.P. and Eibl, J.D.
Improved recombinant eukaryotic cytoplasmic viruses, method for their production and their use as vaccines
Patent: EP 0753581-A 23 15-JAN-1997;
1 (bases 1 to 45)
Scheiflinger, F.D., Antoine, G.D., Falkner, Falko-Guenter, D.,
Scheiflinger, F.D., Antoine, G.D., Falkner, Falko-Guenter, D.,
Improved recombinant eukaryotic cytoplasmic viruses, method their production and their use as vaccines
Patent: EP 0753581-A 24 15-JAN-1997;
                                                                                                unidentified
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Sequence 23 from Patent
A59035
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Method of modulating metabolite biosynthesis in recombinant cells
Patent: Wo 0046383-A 7 10-AUG-2000;
UNIV LEIDEN (NL); MEMELINK JOHAN (NL); FITS CORNELIA THEODORA
ELISABE (NL); KIJNE JAN WILLEM (NL); MENKE FRANK LEONARDUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         unidentified
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/db_xref="taxon:4058"
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                                Au,K.G., Chen,J.G., Patil,N. and Thomas, Genetic compositions and methods Patent: WO 0134840-A 20 17-MAY-2001; GLAXO GROUP LIMITED (GB); Affymetrix, 1 Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Mammalla; Eutheria; Primates; 1 (bases 1 to 19)
Robbins,J.M. and Tritz,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AX130478 19 L
Sequence 1696 from
AX130478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IMMUNO AG
                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                              human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AX130478.1
                                                                                             Mammalia; Eutheria;
1 (bases 1 to 21)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent: WO 0130362-A 1696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                  Conservative
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
/note="Cyclin C ribozyme binding
2 c 4 g 5 t
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/db_xref="taxon:32644"
14 c 12 g 1
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                                                                                                                                                                                                                                                                                                                              55.0%;
83.3%;
                                                                                                                                                                     GI:14284347
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93.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent
                                                                                                       Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                Score 13.2; DB 6;
Pred. No. 4.5e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 13.4; DB 6; Pred. No. 3.7e+04;
                                                                                                                                                                                             WO0134840.
                                                                                                                                                                                                        DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-MAY-2001;
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                                                                                                       Craniata; Ve
Catarrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11
                                               Inc.
                                                                                                                     Vertebrata; Euteleostomi;
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                                                                                                                                                                                                                                                                                                                                                                                                    site"
                                                                                                          Hominidae;
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                                             (US)
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AUTHORS
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ORIGIN
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AX111191
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ORIGIN
Search completed: March 9, 2002, 00:48:44
Job time: 11125 sec
                                                                                                                                                                                                                                                                                                                                                   FEATURES
                                                                                                                                                     Query Match 55.0%; Score 13.2; DB 6; Length 23; Best Local Similarity 83.3%; Pred. No. 4.5e+04; Matches 15; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ouery Match 55.0%; Score 13.2; DB 6; Length 21; Best Local Similarity 78.9%; Pred. No. 4.5e+04; Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE
                                                                                                                                                                                                                                                                                                                                                                                     JOURNAL
                                                                                                                                                                                                                                                                                                                                  source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      variation
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||||| || ||||| |||
5 TCCAATGCTGAGCACTTT 22
                                                                                                                                                                                                                                                                                                                           synthetic construct.

synthetic construct
artificial sequence.

1 (bases 1 to 23)

Bergeron,M.G., Boissinot,M., Huletsky,A., m Nard,C., Ouellette,M.,
Picard,F.J. and Roy,P.H.

Highly conserved genes and their use to generate probes and primers
for detection of microorganisms
Patent: WO 0123604-A 1924 05-APR-2001;
Infectio Diagnostic (I.D.I.) INC. (CA)
Location/Qualifiers

1. 23
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AX111191 23 bp DNA
Sequence 1924 from Patent WO0123604.
AX111191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AX111191.1 GI:13927483
                                                                                                                                                                                                                                                         /organism="synthetic construct"
/db_xref="taxon:32630"
/note="Oligonucleotide"
10 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="'n' represents a polymorphic base" 4 c 	 4 g 	 6 t 	 1 	 others
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